

Title: Autonomous Data Mining.
Inventor: Andrew CONWAY
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Technical Appendix

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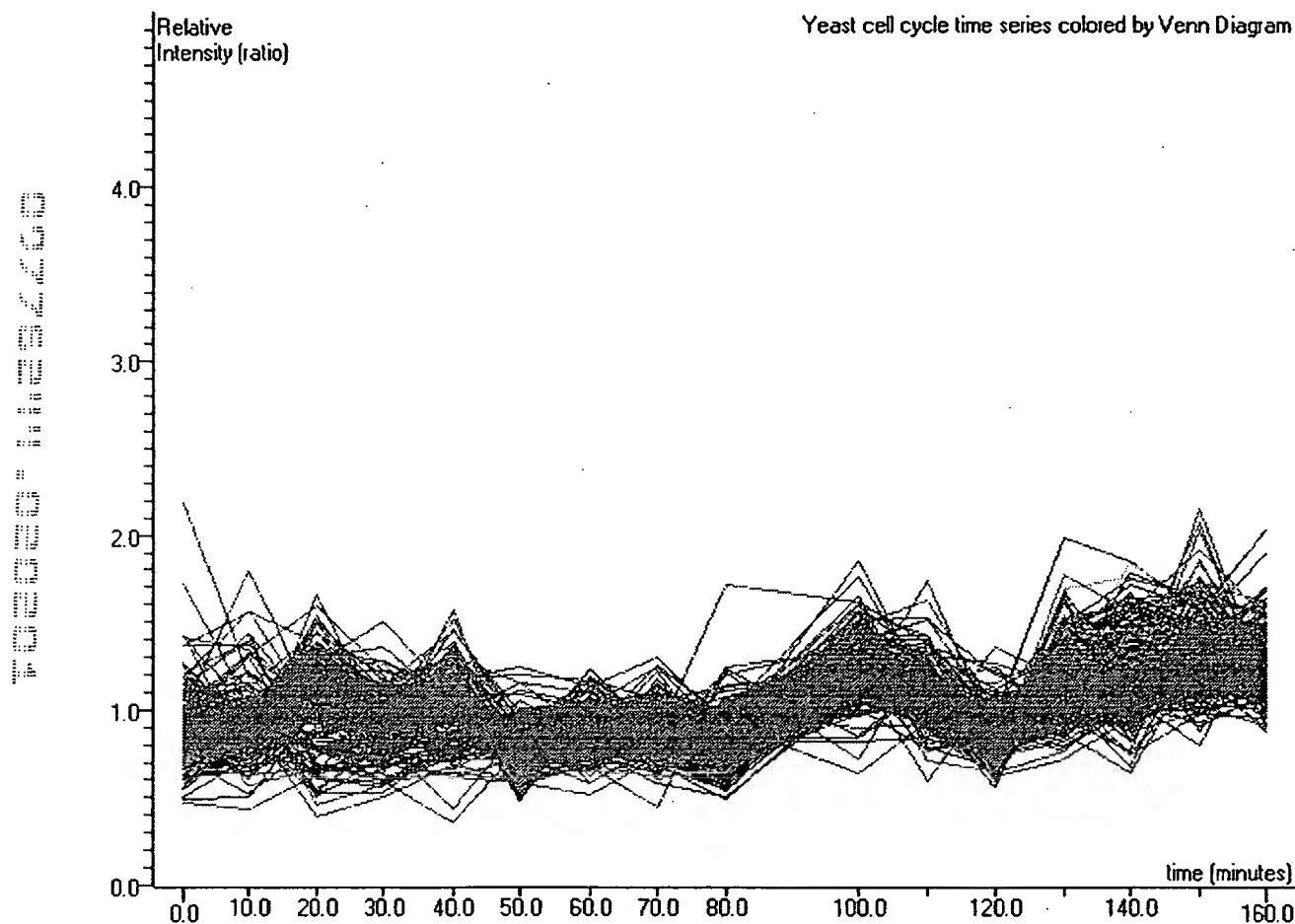
Yeast cell cycle time series and ribosome

I was looking through your experiment called 'Yeast cell cycle time series' and found something I think you will find interesting.

I have found a set of 67 genes with similar expression patterns in 'Yeast cell cycle time series'. These genes are statistically similar to genes in another gene list containing 159 genes called 'ribosome'. 'ribosome' is a Gene List in the folder 'PIR keywords'. It was made by Andrew Conway from Silicon Genetics. These two lists share 18 genes in common.

There is of course a chance this is just a coincidence. I rate that chance at one in 1,252,202,449.

A picture is attached. If you would like to look at this in greater detail, please use the attached GeneSpring bookmark, [Report23haystack.gsp](#) or follow this link: [GenEx](#)



The co-regulated genes are plotted in red, and the genes in ribosome are plotted in green. If a gene is both, it is colored yellow.

Genes in both lists:

YBR048W (RPS11B)

YDL081C (RPP1A)

YDL130W (RPP1B)

YDR447C (RPS17B)

YDR450W (RPS18A)

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YDR471W (RPL27B)

YDL136W (RPL35B)

YDL082W (RPL13A)

Higa S, Yoshihama M, Tanaka T, Kenmochi N. Gene organization and sequence of the region containing the ribosomal protein genes RPL13A and RPS11 in the human genome and conserved features in the mouse genome. Gene. 1999 Nov 29;240(2):371-7.

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YDL083C (RPS16B)

YDR012W (RPL4B)

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YDL061C (RPS29B)

YDL075W (RPL31A)

YDR382W (RPP2B)

YCR031C (RPS14A)

YDR500C (RPL37B)

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YGL030W (RPL30)

Li B, Nierras CR, Warner JR. Transcriptional elements involved in the repression of ribosomal protein synthesis. Mol Cell Biol. 1999 Aug;19(8):5393-404.

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YIL148W (RPL40A)

Coregulated genes not in ribosome

YAL003W (EFB1)

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YAL038W (CDC19)

Ogawa Y, Takahashi T, Masukata H. Association of fission yeast Orp1 and Mcm6 proteins with chromosomal replication origins. Mol Cell Biol. 1999 Oct;19(10):7228-36.

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Barendse W, Armitage SM, Aleyasin A, Womack JE. Differences between the radiation hybrid and genetic linkage maps of bovine chromosome 5 resolved with a quasi-phylogenetic method of analysis. Mamm Genome. 2000 May;11(5):369-72.

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YDR025W (RPS11A)

YDL208W (NHP2)

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